# **WEST Search History**



DATE: Thursday, April 22, 2004

Hide?	Hit Count			
	DB = USF	PT; PLUR=YES; OP=ADJ		
	L8	13 and optimal grow\$	8	
	L7	13 and optimal temperature	4	
	L6	13 and temperature	45	
	L5	L3 and osmoti\$	, 6	
	L4	13 and viral vector	11	
	L3	L2 and plant	45	
	L2	L1 and complement?	56	
	L1	conditional lethal	223	

**END OF SEARCH HISTORY** 

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NEWS 4 JAN 27 A new search aid, the Company Name Thesaurus, available in CA/CAplus

NEWS 5 FEB 05 German (DE) application and patent publication number format changes

NEWS 6 MAR 03 MEDLINE and LMEDLINE reloaded

NEWS 7 MAR 03 MEDLINE file segment of TOXCENTER reloaded

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NEWS 12 MAR 29 New monthly current-awareness alert (SDI) frequency in RAPRA

NEWS EXPRESS MARCH 31 CURRENT WINDOWS VERSION IS V7.00A, CURRENT MACINTOSH VERSION IS V6.0c(ENG) AND V6.0Jc(JP), AND CURRENT DISCOVER FILE IS DATED 13 APRIL 2004

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=> FIL STNGUIDE COST IN U.S. DOLLARS

SINCE FILE TOTAL ENTRY SESSION 0.21 0.21

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COST IN U.S. DOLLARS SINCE FILE TOTAL
ENTRY SESSION
FULL ESTIMATED COST 0.06 0.27

FILE 'HOME' ENTERED AT 10:41:42 ON 22 APR 2004

=> file agricola caplus biosis

COST IN U.S. DOLLARS
SINCE FILE TOTAL
ENTRY SESSION
FULL ESTIMATED COST
0.21
0.48

FILE 'AGRICOLA' ENTERED AT 10:41:50 ON 22 APR 2004

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FILE 'BIOSIS' ENTERED AT 10:41:50 ON 22 APR 2004 COPYRIGHT (C) 2004 BIOLOGICAL ABSTRACTS INC.(R)

=> s conditional lethal

L1 1029 CONDITIONAL LETHAL

=> s l1 and complement?

L2 176 L1 AND COMPLEMENT?

=> s 12 and plant?

L3 38 L2 AND PLANT?

=> dup rem 13

PROCESSING COMPLETED FOR L3

L4 33 DUP REM L3 (5 DUPLICATES REMOVED)

=> d 1-33 ti

- L4 ANSWER 1 OF 33 BIOSIS COPYRIGHT 2004 BIOLOGICAL ABSTRACTS INC. on STN TI The biophysical characterization of putative VDAC isoforms from Drosophila melanogaster.
- L4 ANSWER 2 OF 33 BIOSIS COPYRIGHT 2004 BIOLOGICAL ABSTRACTS INC. on STN TI Poly(A) tail-dependent exonuclease AtRrp41p from Arabidopsis thaliana
- Poly(A) tail-dependent exonuclease AtRrp41p from Arabidopsis thaliana rescues 5.8 S rRNA processing and mRNA decay defects of the yeast ski6 mutant and is found in an exosome-sized complex in plant and yeast cells.
- L4 ANSWER 3 OF 33 BIOSIS COPYRIGHT 2004 BIOLOGICAL ABSTRACTS INC. on STN
- TI Cloning and characterization of the Schizosaccharomyces pombe tRNA:pseudouridine synthase Puslp.
- L4 ANSWER 4 OF 33 BIOSIS COPYRIGHT 2004 BIOLOGICAL ABSTRACTS INC. on STN
- TI An additional transcript of the cdc25C gene from A431 cells encodes a functional protein.
- L4 ANSWER 5 OF 33 BIOSIS COPYRIGHT 2004 BIOLOGICAL ABSTRACTS INC. on STN
- TI Cloning and sequencing of the Candida albicans C-4 sterol methyl oxidase gene (ERG25) and expression of an ERG25 conditional lethal mutation in Saccharomyces cerevisiae.
- L4 ANSWER 6 OF 33 BIOSIS COPYRIGHT 2004 BIOLOGICAL ABSTRACTS INC. on STN

- TI Chromosomal promoter replacement in Saccharomyces cerevisiae: Construction of conditional lethal strains for the cloning of glycosyltransferases from various organisms.
- L4 ANSWER 7 OF 33 AGRICOLA Compiled and distributed by the National Agricultural Library of the Department of Agriculture of the United States of America. It contains copyrighted materials. All rights reserved.

  (2004) on STN
- TI A tomato homeobox gene (HD-Zip) is involved in limiting the spread of programmed cell death.
- L4 ANSWER 8 OF 33 BIOSIS COPYRIGHT 2004 BIOLOGICAL ABSTRACTS INC. on STN
- TI Deletion analysis of yeast Sec65p reveals a central domain that is sufficient for function in vivo.
- L4 ANSWER 9 OF 33 BIOSIS COPYRIGHT 2004 BIOLOGICAL ABSTRACTS INC. on STN
- Cloning and functional expression of the human GlcNAc-1-P transferase, the enzyme for the committed step of the dolichol cycle, by heterologous complementaion in Saccharomyces cerevisiae.
- L4 ANSWER 10 OF 33 AGRICOLA Compiled and distributed by the National Agricultural Library of the Department of Agriculture of the United States of America. It contains copyrighted materials. All rights reserved.

  (2004) on STN DUPLICATE 1
- TI Molecular cloning and functional analysis of the Arabidopsis thaliana DNA ligase I homologue.
- L4 ANSWER 11 OF 33 BIOSIS COPYRIGHT 2004 BIOLOGICAL ABSTRACTS INC. on STN
- TI Casein kinase II regulation of yeast TFIIIB is mediated by the TATA-binding protein.
- L4 ANSWER 12 OF 33 BIOSIS COPYRIGHT 2004 BIOLOGICAL ABSTRACTS INC. on STN
- TI BIM1 encodes a microtubule-binding protein in yeast.
- L4 ANSWER 13 OF 33 BIOSIS COPYRIGHT 2004 BIOLOGICAL ABSTRACTS INC. on STN
- TI A conditional-lethal translation termination defect in a sup45 mutant of the yeast Saccharomyces cerevisiae.
- L4 ANSWER 14 OF 33 AGRICOLA Compiled and distributed by the National Agricultural Library of the Department of Agriculture of the United States of America. It contains copyrighted materials. All rights reserved.

  (2004) on STN DUPLICATE 2
- TI Evidence for multiple forms of biotin holocarboxylase synthetase in pea (Pisum sativum) and in Arabidopsis thaliana: subcellular fractionation studies and isolation of a cDNA clone.
- L4 ANSWER 15 OF 33 BIOSIS COPYRIGHT 2004 BIOLOGICAL ABSTRACTS INC. on STN
- TI Slslp, an endoplasmic reticulum component, is involved in the protein translocation process in the yeast Yarrowia lipolytica.
- L4 ANSWER 16 OF 33 BIOSIS COPYRIGHT 2004 BIOLOGICAL ABSTRACTS INC. on STN
- TI Human ARF4 expression rescues sec7 mutant yeast cells.
- L4 ANSWER 17 OF 33 BIOSIS COPYRIGHT 2004 BIOLOGICAL ABSTRACTS INC. on STN
- TI RNA polymerase III defects suppress a conditional-lethal poly(A) polymerase mutation in Saccharomyces cerevisiae.
- L4 ANSWER 18 OF 33 BIOSIS COPYRIGHT 2004 BIOLOGICAL ABSTRACTS INC. on STN
- TI A connection between pre-mRNA splicing and the cell cycle in fission yeast: cdc28+ is allelic with prp8+ and encodes an RNA-dependent ATPase/helicase.
- L4 ANSWER 19 OF 33 BIOSIS COPYRIGHT 2004 BIOLOGICAL ABSTRACTS INC. on STN
- TI Structure-based systematic isolation of conditional-

lethal mutations in the single yeast calmodulin gene.

- L4 ANSWER 20 OF 33 BIOSIS COPYRIGHT 2004 BIOLOGICAL ABSTRACTS INC. on STN
- TI The identification of a Caenorhabditis elegans homolog of p34-cdc2 kinase.
- L4 ANSWER 21 OF 33 BIOSIS COPYRIGHT 2004 BIOLOGICAL ABSTRACTS INC. on STN
- TI Cloning of a gene (PSD1) encoding phosphatidylserine decarboxylase from Saccharomyces cerevisiae by complementation of an Escherichia coli mutant.
- L4 ANSWER 22 OF 33 BIOSIS COPYRIGHT 2004 BIOLOGICAL ABSTRACTS INC. on STN
- TI Fission yeast with DNA polymerase delta temperature-sensitive alleles exhibits cell division cycle phenotype.
- L4 ANSWER 23 OF 33 BIOSIS COPYRIGHT 2004 BIOLOGICAL ABSTRACTS INC. on STN
- TI In vivo species specificity of DNA polymerase alpha.
- L4 ANSWER 24 OF 33 BIOSIS COPYRIGHT 2004 BIOLOGICAL ABSTRACTS INC. on STN
- TI THE YEAST SLY GENE PRODUCTS SUPPRESSORS OF DEFECTS IN THE ESSENTIAL GTP-BINDING YPT1 PROTEIN MAY ACT IN ENDOPLASMIC RETICULUM-TO-GOLGI TRANSPORT.
- L4 ANSWER 25 OF 33 BIOSIS COPYRIGHT 2004 BIOLOGICAL ABSTRACTS INC. on STN
- TI EXPRESSION OF HUMAN DNA TOPOISOMERASE I IN YEAST CELLS LACKING YEAST DNA TOPOISOMERASE I RESTORATION OF SENSITIVITY OF THE CELLS TO THE ANTITUMOR DRUG CAMPTOTHECIN.
- L4 ANSWER 26 OF 33 BIOSIS COPYRIGHT 2004 BIOLOGICAL ABSTRACTS INC. on STN
- TI A YEAST GENE IMPORTANT FOR PROTEIN ASSEMBLY INTO THE ENDOPLASMIC RETICULUM AND THE NUCLEUS HAS HOMOLOGY TO DNAJ AN ESCHERICHIA-COLI HEAT SHOCK PROTEIN.
- L4 ANSWER 27 OF 33 BIOSIS COPYRIGHT 2004 BIOLOGICAL ABSTRACTS INC. on STN
- TI FUNCTIONAL EXPRESSION OF CHICKEN CALMODULIN IN YEAST.
- L4 ANSWER 28 OF 33 CAPLUS COPYRIGHT 2004 ACS on STN DUPLICATE 3
- TI Biochemical genetics of further chlorate resistant molybdenum cofactor defective, conditional-lethal mutants of barley
- L4 ANSWER 29 OF 33 BIOSIS COPYRIGHT 2004 BIOLOGICAL ABSTRACTS INC. on STN
- TI PROTEIN SORTING IN SACCHAROMYCES-CEREVISIAE ISOLATION OF MUTANTS DEFECTIVE IN THE DELIVERY AND PROCESSING OF MULTIPLE VACUOLAR HYDROLASES.
- L4 ANSWER 30 OF 33 BIOSIS COPYRIGHT 2004 BIOLOGICAL ABSTRACTS INC. on STN
- TI A COMPLEMENTATION ANALYSIS BY PARASEXUAL RECOMBINATION OF CANDIDA-ALBICANS MORPHOLOGICAL MUTANTS.
- L4 ANSWER 31 OF 33 BIOSIS COPYRIGHT 2004 BIOLOGICAL ABSTRACTS INC. on STN
- TI THE ALLOSUPPRESSOR GENE SAL-4 ENCODES A PROTEIN IMPORTANT FOR MAINTAINING TRANSLATIONAL FIDELITY IN SACCHAROMYCES-CEREVISIAE.
- L4 ANSWER 32 OF 33 BIOSIS COPYRIGHT 2004 BIOLOGICAL ABSTRACTS INC. on STN
- TI UNLINKED NONCOMPLEMENTATION ISOLATION OF NEW CONDITIONAL-LETHAL MUTATIONS IN EACH OF THE TUBULIN GENES OF SACCHAROMYCES-CEREVISIAE.
- L4 ANSWER 33 OF 33 BIOSIS COPYRIGHT 2004 BIOLOGICAL ABSTRACTS INC. on STN
- TI ISOLATION OF THE BETA TUBULIN GENE FROM YEAST SACCHAROMYCES-CEREVISIAE AND DEMONSTRATION OF ITS ESSENTIAL FUNCTION IN-VIVO.

=> d 7 ab

L4 ANSWER 7 OF 33 AGRICOLA Compiled and distributed by the National

Agricultural Library of the Department of Agriculture of the United States of America. It contains copyrighted materials. All rights reserved. (2004) on STN

Antisense suppression in transgenic tomato plants of H52, a gene encoding a new homeodomain protein of the HD-Zip class, produces a conditional lethal phenotype. The transgenic lines that survive exhibit spontaneous misregulation of cell death control in leaves, which, once initiated, propagates and engulfs the entire leaf. Activation of defence genes, over-accumulation of ethylene and conjugated salicylic acid, and growth reduction of virulent pathogens also occurs in these plants. In wild-type plants, H52 is up-regulated upon infection, mirroring the generation of the oxidative burst which normally precedes the hypersensitive response (HR). Thus, H52 appears to be a transcription factor involved in cellular protection by limiting spread of programmed cell death in plants.

#### => d 7 so 9

'SO9' IS NOT A VALID FORMAT

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REENTER DISPLAY FORMAT FOR ALL FILES (FILEDEFAULT):d 7 so

'D' IS NOT A VALID FORMAT

'7' IS NOT A VALID FORMAT

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REENTER DISPLAY FORMAT FOR ALL FILES (FILEDEFAULT):ti

- L4 ANSWER 7 OF 33 AGRICOLA Compiled and distributed by the National Agricultural Library of the Department of Agriculture of the United States of America. It contains copyrighted materials. All rights reserved. (2004) on STN
- TI A tomato homeobox gene (HD-Zip) is involved in limiting the spread of programmed cell death.

# => d 7 so

- L4 ANSWER 7 OF 33 AGRICOLA Compiled and distributed by the National Agricultural Library of the Department of Agriculture of the United States of America. It contains copyrighted materials. All rights reserved. (2004) on STN
- SO The Plant journal : for cell and molecular biology, Dec 1999. Vol. 20, No. 5. p. 591-600
  Publisher: Oxford : Blackwell Sciences Ltd.
  ISSN: 0960-7412

# => d 10 ab

- L4 ANSWER 10 OF 33 AGRICOLA Compiled and distributed by the National Agricultural Library of the Department of Agriculture of the United States of America. It contains copyrighted materials. All rights reserved.

  (2004) on STN DUPLICATE 1
- AB A cDNA encoding the DNA ligase I homologue has been isolated from Arabidopsis thaliana using a degenerate PCR approach. The ORF of this cDNA encodes an amino acid sequence of 790 residues, representing a protein with a theoretical molecular mass of 87.8 kDa and an isoelectric point (pl) of 8.20. Alignment of the A. thaliana DNA ligase protein sequence with the sequence of DNA ligases from human (Homo sapiens), murine (Mus

musculus), clawed toad (Xenopus laevis) and the yeasts Schizosaccharomyces pombe and Saccharomyces cerevisiae showed good sequence homology (42-45% identity, 61-66% similarity), particularly around the active site. Sequence data indicate that the Arabidopsis DNA ligase is the homologue of the animal DNA ligase I species. Functional analysis of the cDNA clone demonstrated its ability to complement the conditional lethal phenotype of an S. cerevisiae cdc9 mutant defective in DNA ligase activity, confirming that the cloned sequence encodes an active DNA ligase. The level of the DNA ligase transcript was not increased in A. thaliana seedlings in response to DNA damage induced by a period of enhanced UV-B irradiation. However, the cellular level of the DNA ligase mRNA transcript does correlate with the replicative state of plant cells.

#### => d 10 so

- L4 ANSWER 10 OF 33 AGRICOLA Compiled and distributed by the National Agricultural Library of the Department of Agriculture of the United States of America. It contains copyrighted materials. All rights reserved.

  (2004) on STN DUPLICATE 1
- SO The Plant journal : for cell and molecular biology, Apr 1998. Vol. 14, No. 1. p. 75-81
  Publisher: Oxford : Blackwell Sciences Ltd.
  ISSN: 0960-7412

## => d 28 ab

L4ANSWER 28 OF 33 CAPLUS COPYRIGHT 2004 ACS on STN DUPLICATE 3 Three plants, R9201 and R11301 (from cv. Maris Mink) and R12202 AB (from cv. Golden Promise), were selected by screening M2 populations of barley (Hordeum vulgare) seedlings (mutagenized with azide in the M1) for resistance to 10 mM potassium chlorate. Selections R9201 and R11301 were crossed with the wild-type cv. Maris Mink and anal. of the F2 progeny showed that one quarter lacked shoot nitrate reductase activity. These F2 plants also withered and died in the continuous presence of nitrate as sole nitrogen source. Loss of nitrate reductase activity, withering and death were due in each case to a recessive mutation in a single nuclear gene. All F1 progeny derived from selfing selection R12202 lacked shoot nitrate reductase activity, withered and subsequently died when maintained in the continuous presence of nitrate as sole nitrogen source. All homozygous mutant plants lacked not only shoot nitrate reductase activity but also shoot xanthine dehydrogenase activity. The plants took up nitrate, and possessed wild-type or higher levels of shoot nitrite reductase activity and NADH-cytochrome c reductase activity when treated with nitrate for 18 h. Apparently, loss of shoot nitrate reductase activity, xanthine dehydrogenase activity and withering and death, in the 3 mutants R9201, R11301 and R12202 is due to a mutation affecting the formation of a functional molybdenum cofactor. The mutants possessed wild-type levels of molybdenum. Growth in the presence of unphysiol. high levels of molybdate did not restore shoot nitrate reductase or xanthine dehydrogenase activity. The shoot molybdenum cofactor of R9201 and R12202 is unable to reconstitute NADPH nitrate reductase activity from exts. of the Neurospora crassa nit-1 mutant and dimerize the nitrate reductase subunits present in the resp. barley mutant. The shoot molybdenum cofactor of R11301 is able to effect dimerization of the R11301 nitrate reductase subunits and can reconstitute NADPH-nitrate reductase activity up to 40% of the wild-type molybdenum cofactor levels. The molybdenum cofactor of the roots of R9201 and R11301 is also defective. Genetic anal. demonstrated that R9201, but not R11301, is allelic to R9401 and Az34 (nar-2a), 2 mutants previously shown to be defective in synthesis of molybdenum cofactor. The mutations in R9401 and R9201 gave partial complementation of the nar-2a gene such that

heterozygotes had higher levels of extractable nitrate reductase activity than the homozygous mutants. Thus, (a) the nar-2 gene locus encodes a step in molybdopterin biosynthesis; (b) the mutant R11301 represents a further locus involved in the synthesis of a functional molybdenum cofactor; (c) mutant R12202 is also defective in molybdopterin biosynthesis; and (d) the nar-2 gene locus and the gene locus defined by R11301 govern molybdenum cofactor biosynthesis in both shoot and root.

ANSWER 28 OF 33 CAPLUS COPYRIGHT 2004 ACS on STN DUPLICATE 3 L4 Molecular and General Genetics (1989), 219(3), 421-8 CODEN: MGGEAE; ISSN: 0026-8925 => s ((pogue g?) or (pogue, g?))/au 174 ((POGUE G?) OR (POGUE, G?))/AU => s 15 and conditional lethal 0 L5 AND CONDITIONAL LETHAL => s 15 and complement? 14 L5 AND COMPLEMENT? L7=> dup rem 17 PROCESSING COMPLETED FOR L7 10 DUP REM L7 (4 DUPLICATES REMOVED) => s 18 and plant? L9 6 L8 AND PLANT? => d 1-6 ti

=> d 28 so

- L9 ANSWER 1 OF 6 AGRICOLA Compiled and distributed by the National Agricultural Library of the Department of Agriculture of the United States of America. It contains copyrighted materials. All rights reserved. (2004) on STN
- TI Barley stripe mosaic virus-induced gene silencing in a monocot plant.
- L9 ANSWER 2 OF 6 AGRICOLA Compiled and distributed by the National Agricultural Library of the Department of Agriculture of the United States of America. It contains copyrighted materials. All rights reserved. (2004) on STN
- TI The requirement for a 5' stem-loop structure in brome mosaic virus replication supports a new model for viral positive-strand RNA initiation.
- L9 ANSWER 3 OF 6 CAPLUS COPYRIGHT 2004 ACS on STN
- TI Development of a **plant** viral-vector-based gene expression assay for the screening of yeast cytochrome P450 monooxygenases
- L9 ANSWER 4 OF 6 CAPLUS COPYRIGHT 2004 ACS on STN
- TI Trans-complementing multiple component RNA virus vectors for the delivery and expression of foreign genes
- L9 ANSWER 5 OF 6 CAPLUS COPYRIGHT 2004 ACS on STN
- TI Method of determining the function of nucleotide sequences and the proteins they encode by transfecting the same into a host
- L9 ANSWER 6 OF 6 CAPLUS COPYRIGHT 2004 ACS on STN
- TI Point mutations in the ICR2 motif of brome mosaic virus RNAs debilitate (+)-strand replication

L9 ANSWER 5 OF 6 CAPLUS COPYRIGHT 2004 ACS on STN

The present invention provides methods for rapidly determining the function of nucleic acid sequences by transfecting the same into a host organism to effect expression. Phenotypic and biochem. changes produced thereby are then analyzed to ascertain the function of the nucleic acids which have been transfected into the host organism. The invention also provides methods for silencing endogenous genes by transfecting hosts with nucleic acid sequences to effect expression of the same. The present invention also provides methods for selecting desired functions of RNAs and proteins by the use of virus vectors to express libraries of nucleic acid sequence variants. Moreover, the present invention provides methods for inhibiting an endogenous protease of a plant host.

### => d 5 pi

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ANSWER 5 OF 6 CAPLUS COPYRIGHT 2004 ACS on STN
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    PATENT NO. KIND DATE
                               APPLICATION NO. DATE
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                    A2
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                                       WO 1999-US1164 19990115
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    WO 9936516
                    A3 20000316
    WO 9936516
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            DK, EE, ES, FI, GB, GD, GE, GH, GM, HR, HU, ID, IL, IN, IS, JP,
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            MW, MX, NO, NZ, PL, PT, RO, RU, SD, SE, SG, SI, SK, SL, TJ, TM,
            TR, TT, UA, UG, US, UZ, VN, YU, ZW, AM, AZ, BY, KG, KZ, MD, RU,
            TJ, TM
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    US 2003167512
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                                       US 2002-236508
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5 DUP REM L13 (1 DUPLICATE REMOVED)

#### => d 1-5 ti

- L14 ANSWER 1 OF 5 BIOSIS COPYRIGHT 2004 BIOLOGICAL ABSTRACTS INC. on STN TI Method of compiling a functional gene profile in a **plant** by
  - transfecting a nucleic acid sequence of a donor plant into a different host plant in an anti-sense orientation.
- L14 ANSWER 2 OF 5 AGRICOLA Compiled and distributed by the National Agricultural Library of the Department of Agriculture of the United States of America. It contains copyrighted materials. All rights reserved.

  (2004) on STN
- TI Rapid, high-level expression of glycosylated rice alpha-amylase in transfected **plants** by an RNA viral vector.
- L14 ANSWER 3 OF 5 CAPLUS COPYRIGHT 2004 ACS on STN
- TI Method of determining the function of nucleotide sequences and the proteins they encode by transfecting the same into a host
- L14 ANSWER 4 OF 5 AGRICOLA Compiled and distributed by the National Agricultural Library of the Department of Agriculture of the United States of America. It contains copyrighted materials. All rights reserved.

  (2004) on STN DUPLICATE 1
- TI Cytoplasmic inhibition of carotenoid biosynthesis with virus-derived RNA.
- L14 ANSWER 5 OF 5 BIOSIS COPYRIGHT 2004 BIOLOGICAL ABSTRACTS INC. on STN
- TI TRANSLOCATION OF THE PRECURSOR OF 5 ENOLPYRUVYLSHIKIMATE-3-PHOSPHATE SYNTHASE INTO CHLOROPLASTS OF HIGHER **PLANTS** IN-VITRO.

#### => d so

L14 ANSWER 1 OF 5 BIOSIS COPYRIGHT 2004 BIOLOGICAL ABSTRACTS INC. on STN Official Gazette of the United States Patent and Trademark Office Patents, (July 30, 2002) Vol. 1260, No. 5. http://www.uspto.gov/web/menu/patdata.html. e-file.

CODEN: OGUPE7. ISSN: 0098-1133.

## => d pi

L14 ANSWER 1 OF 5 BIOSIS COPYRIGHT 2004 BIOLOGICAL ABSTRACTS INC. on STN PI US 6426185 July 30, 2002

# => d 3 pi

L	14	ANSWER PATENT		_			COPY!							٥.	DATE			
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            419 ((ERWIN, R?) OR (ERWIN R?))/AU
=> s 115 and conditional lethal
               0 L15 AND CONDITIONAL LETHAL
=> s l15 and complement?
              4 L15 AND COMPLEMENT?
=> dup rem 117
PROCESSING COMPLETED FOR L17
                4 DUP REM L17 (0 DUPLICATES REMOVED)
=> d 1-4 ti
L18 ANSWER 1 OF 4 BIOSIS COPYRIGHT 2004 BIOLOGICAL ABSTRACTS INC. on STN
     Method of compiling a functional gene profile in a plant by transfecting a
     nucleic acid sequence of a donor plant into a different host plant in an
     anti-sense orientation.
L18 ANSWER 2 OF 4 CAPLUS COPYRIGHT 2004 ACS on STN
     Method of determining the function of nucleotide sequences and the
     proteins they encode by transfecting the same into a host
L18 ANSWER 3 OF 4 BIOSIS COPYRIGHT 2004 BIOLOGICAL ABSTRACTS INC. on STN
     MIDLATENCY AUDITORY EVOKED RESPONSES P1 ABNORMALITIES IN ADULT AUTISTIC
     SUBJECTS.
L18 ANSWER 4 OF 4 BIOSIS COPYRIGHT 2004 BIOLOGICAL ABSTRACTS INC. on STN
     RELATIONS AMONG CLINICAL SCALES IN SCHIZOPHRENIA.
=> s ((hanley k?) or (hanley, k?))/au
            237 ((HANLEY K?) OR (HANLEY, K?))/AU
=> s 119 and conditional lethal
              0 L19 AND CONDITIONAL LETHAL
=> s 119 and complement?
            10 L19 AND COMPLEMENT?
=> s 121 and plant?
              8 L21 AND PLANT?
=> dup rem 122
PROCESSING COMPLETED FOR L22
               5 DUP REM L22 (3 DUPLICATES REMOVED)
```

T-15

L17

TТ

TΤ

L22

=> d 1-5 ti

L23 ANSWER 1 OF 5 CAPLUS COPYRIGHT 2004 ACS on STN

- TI Development of a **plant** viral-vector-based gene expression assay for the screening of yeast cytochrome P450 monooxygenases
- L23 ANSWER 2 OF 5 CAPLUS COPYRIGHT 2004 ACS on STN
- TI Method of determining the function of nucleotide sequences and the proteins they encode by transfecting the same into a host
- L23 ANSWER 3 OF 5 CAPLUS COPYRIGHT 2004 ACS on STN
- TI Squalene synthetase cDNA sequence of Nicotiana benthamiana for sterol formation
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  (2004) on STN

  DUPLICATE 1
- TI Molecular cloning, in vitro expression and characterization of a plant squalene synthetase cDNA.
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  (2004) on STN

  DUPLICATE 2
- TI Cytoplasmic inhibition of carotenoid biosynthesis with virus-derived RNA.

#### => d 3 ab

- L23 ANSWER 3 OF 5 CAPLUS COPYRIGHT 2004 ACS on STN
- AB A DNA sequence isolated as a cDNA from a Nicotiana species, e.g., Nicotiana benthamiana, has a nucleotide sequence which encodes a native squalene synthetase capable of conducting the reductive condensation of two mols. of farnesyl diphosphate to form squalene, constituting the first committed step in sterol biosynthesis in eukaryotes.

# => d 3 so

L23 ANSWER 3 OF 5 CAPLUS COPYRIGHT 2004 ACS on STN SO PCT Int. Appl., 46 pp.
CODEN: PIXXD2

## => d 3 pi

L23 ANSWER 3 OF 5 CAPLUS COPYRIGHT 2004 ACS on STN PATENT NO. KIND DATE APPLICATION NO. DATE ---- ------\_\_\_\_\_\_ A1 19960328 PΤ WO 9609393 WO 1995-US11280 19950907 W: AM, AT, AU, BB, BG, BR, BY, CA, CH, CN, CZ, DE, DK, EE, ES, FI, GB, GE, HU, IS, JP, KE, KG, KP, KR, KZ, LK, LR, LT, LU, LV, MD, MG, MK, MN, MW, MX, NO, NZ, PL, PT, RO, RU, SD, SE, SG, SI, SK, TJ, TM RW: KE, MW, SD, SZ, UG, AT, BE, CH, DE, DK, ES, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE, BF, BJ, CF, CG, CI, CM, GA, GN, ML, MR, NE, SN, TD, TG US 5741898 Α 19980421 US 1994-310693 19940922 AU 9535465 A1 19960409 AU 1995-35465 19950907

## => d 4 ab

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(2004) on STN DUPLICATE 1

AB

Squalene synthetase (farnesyl-diphosphate:farnesyl-diphosphate farnesyltransferase, EC 2.5.1.21) catalyzes the first committed step for sterol biosynthesis and is thought to play an important role in the regulation of isoprenoid biosynthesis in eukaryotes. Using degenerate oligonucleotides based on a conserved region found in yeast and human squalene synthetase genes, a cDNA was cloned from the plant Nicotiana benthamiana. The cloned cDNA contained an open reading frame of 1234 bp encoding a polypeptide of 411 amino acids (Mr 47002). Northern blot analysis of poly(A) + mRNA from N. benthamiana and N. tabacum cv. MD609 revealed a single band of ca. 1.6 kb in both Nicotiana species. The identity and functionality of the cloned plant squalene synthetase cDNA was further confirmed by expression of the cDNA in Escherichia coli and in a squalene synthetase-deficient erg9 mutant of Saccharomyces cerevisiae. Antibodies raised against a truncated form of the protein recognized an endogenous plant protein of appropriate size as well as the full-length bacterially expressed protein as detected by western analysis. Comparison of the deduced primary amino acid sequences of plant, yeast, rat and human squalene synthetase revealed regions of conservation that may indicate similar functions within each polypeptide.